



IFWO

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/791,860

DATE: 08/18/2004

TIME: 09:08:21

Input Set : A:\2004-08-12 0641-0260P.ST25.txt
 Output Set: N:\CRF4\08182004\J791860.raw

3 <110> APPLICANT: Bing-Ren HUANG et al.
 5 <120> TITLE OF INVENTION: REGULATOR OF APOPTOSIS AND CELL PROLIFERATION
 7 <130> FILE REFERENCE: 0641-0260P
 9 <140> CURRENT APPLICATION NUMBER: US 10/791,860
 10 <141> CURRENT FILING DATE: 2004-03-04
 12 <160> NUMBER OF SEQ ID NOS: 20
 14 <170> SOFTWARE: PatentIn version 3.2
 16 <210> SEQ ID NO: 1
 17 <211> LENGTH: 197
 18 <212> TYPE: DNA
 19 <213> ORGANISM: Rattus norvegicus
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 28 ttgcgaaaa aaaaaaaaaa 197
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 36 <400> SEQUENCE: 2
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 39 ttccctccctg ctgttggag gcagcatctc ctcttttat ggagggcccg tccttttcc 120
 41 ttacaaattt ttcaataaaag acacattttt gaggcgaaaa aaaaaaaaaa 167
 44 <210> SEQ ID NO: 3
 45 <211> LENGTH: 901
 46 <212> TYPE: DNA
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 54 gactatctca tgagtacgc cttctggggc ccagttccca actgggtct cccattgct 180
 56 gctatcaatg acatgaagaa atctccagag attatcgtt ggcggatgac tttcccccctc 240
 58 tggcttatt ctctgacatt catgagattt gcctacaagg tacaaccccg aaactggctt 300
 60 ctgtttgcgt gccatgtgac aaacgaagtc gctcgttca ttcaggagg acgacttac 360
 62 aactacgaga tgagtaagcg gccatctgcc tagcaatgcg aggaccagct cttgaaagg 420
 64 acagtgcgtc agccactgtt gggccacag atcacgtcgtt catgaatagt cgtgctgagg 480
 66 gaaaaacacg gaagactatc ttaatgacc atgccaacat tattgaatag ccaagaatcc 540
 68 ccaaaaccaac tctcgctgc cttatcaatg ctttttttta tttgtcttca tcaggatgt 600
 70 ttcaaaatattt gcaatgtttaataattttt gaatgtatgtt atctatagca atctgttagta 660
 72 atatgtatattatcttattgg gatttgcgtt ataaaaaaatc taaggaaaca aaactttata 720
 74 actacaagca cttaagtccct caaaatttctt gacttttctt ttaatgacta tagtataacc 780
 76 ctcagttggc cacatgttca cacataattt ccagtgataa caagtagccgg tttttccat 840



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78 atgtaattca gatctgaact taatggcaat aaatggttta aatatttgcg aaaaaaaaaaa 900
 80 a 901
 83 <210> SEQ ID NO: 4
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 85 <212> TYPE: PRT
 86 <213> ORGANISM: Rattus norvegicus
 88 <400> SEQUENCE: 4
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 91 1 5 10 15
 94 Asp Phe Arg Asp Tyr Leu Met Ser Thr His Phe Trp Gly Pro Val Ala
 95 20 25 30
 98 Asn Trp Gly Leu Pro Ile Ala Ala Ile Asn Asp Met Lys Lys Ser Pro
 99 35 40 45
 102 Glu Ile Ile Ser Gly Arg Met Thr Phe Ala Leu Cys Cys Tyr Ser Leu
 103 50 55 60
 106 Thr Phe Met Arg Phe Ala Tyr Lys Val Gln Pro Arg Asn Trp Leu Leu
 107 65 70 75 80
 110 Phe Ala Cys His Val Thr Asn Glu Val Ala Gln Leu Ile Gln Gly Gly
 111 85 90 95
 114 Arg Leu Ile Asn Tyr Glu Met Ser Lys Arg Pro Ser Ala
 115 100 105
 118 <210> SEQ ID NO: 5
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 120 <212> TYPE: PRT
 121 <213> ORGANISM: Mus musculus
 123 <400> SEQUENCE: 5
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 126 1 5 10 15
 129 Asp Phe Arg Asp Tyr Leu Met Ser Thr His Phe Trp Gly Pro Val Ala
 130 20 25 30
 133 Asn Trp Gly Leu Pro Ile Ala Ala Ile Asn Asp Met Lys Lys Ser Pro
 134 35 40 45
 137 Glu Ile Ile Ser Gly Arg Met Thr Phe Ala Leu Cys Cys Tyr Ser Gln
 138 50 55 60
 141 Thr Phe Met Arg Phe Ala Tyr Lys Val Gln Pro Arg Asn Trp Leu Leu
 142 65 70 75 80
 145 Phe Ala Cys His Val Thr Asn Glu Val Ala Gln Leu Ile Gln Gly Gly
 146 85 90 95
 149 Arg Leu Ile Asn Tyr Glu Met Ser Lys Arg Pro Ser Ala
 150 100 105
 153 <210> SEQ ID NO: 6
 154 <211> LENGTH: 102
 155 <212> TYPE: PRT
 156 <213> ORGANISM: Homo sapiens
 158 <400> SEQUENCE: 6
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 161 1 5 10 15
 164 Asp Phe Arg Asp Tyr Leu Met Ser Thr His Phe Trp Gly Pro Val Ala
 165 20 25 30

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168 Asn Trp Gly Leu Pro Ile Ala Ala Ile Asn Asp Met Lys Lys Ser Pro
169          35          40          45
172 Glu Ile Ile Ser Gly Arg Met Thr Phe Ala Leu Cys Cys Tyr Ser Leu
173          50          55          60
176 Thr Phe Met Arg Phe Ala Tyr Lys Val Gln Pro Arg Asn Trp Leu Leu
177 65          70          75          80
180 Phe Ala Cys His Ala Thr Asn Glu Val Ala Gln Leu Ile Gln Gly Gly
181          85          90          95
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185          100
188 <210> SEQ ID NO: 7

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188 <210> SEQ ID NO: 7

189 <211> LENGTH: 9

190 <212> TYPE: DNA

191 <213> ORGANISM:

191 <213> ORGANISM: HOMO sapiens
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193 <400> SEQUENCE: 7
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194	gcgcgtggggc	gggccttcgg	gctggctcg	cgtcggctgc	ccccgggggtt	gcctgggtgt	60
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198	ccatggcggg	cgcgttgggt	cgaaaagcgg	cggactatgt	ccgaagcaag	gattccggg	180
200	actacctcat	gagtacgcac	ttctggggcc	cagtagccaa	ctggggtctt	ccattgtctg	240
202	ccatcaatga	tatgaaaaag	tctccagaga	ttatcagtgg	gcggatgaca	tttgcctct	300
204	gttgctattc	tttgacattc	atgagatttgc	cctacaaggt	acagcctcgg	aactggcttc	360
206	tgtttgcatg	ccacgcaaca	aatgaagtag	cccagctcat	ccagggaggg	cggcttatca	420
208	aacacgagat	gactaaaacg	gcatctgc	aacaatgggaa	aaaggaagaa	caaggtcttgc	480
210	aaggcacgc	attgccagct	gctgctgagt	cacagatttgc	attataaata	gcctccctaa	540
212	ggaaaaataca	ctgaatgcta	ttttactaa	ccattctatt	tttataaaaa	tagtgcggag	600
214	tttctaaacc	aactctctgc	tgccttacaa	gtattaaata	tttacttct	ttccatcaaag	660
216	agtagctcaa	aatatgcaat	taatattaata	atttctgtat	atgttttata	tgcagtaata	720
218	tgtatatacat	ctattagaat	ttacttaatgc	aaaaactgaa	gagaacaaaa	tttgcggacca	780
220	ctagcactta	agtactcctgc	attcttaaca	ttgtctttaa	tgaccacaag	acaaccaaca	840
222	gctggccacgc	tactaaaaat	tttgcggcc	ctgtttaaaa	atgttacctgc	tgtatccat	900
224	tgcagtgtat	atattggat	gctgttaact	aatggcaata	aatgatttaa	atatttgcgtt	960
226	aaaaaaaaaaa	aaaaaaaaaaa	aaaaaaaaaaa				988

229 <210> SEQ ID NO: 8

230 <211> LENGTH: 873

231 <212> TYPE: DNA

232 <213> ORGANISM: *Mus musculus*

234 <400> SEQUENCE: 8

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234 <400> SEQUENCE: 6
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237 gtgcgcaaag cggcggacta tgcggagc aaggacttcc gggactatct catgagtacg 120
239 cacttctggg gcccagggtc caactgggtt ctccccattt ctgttatcaa tgacatgaag 180
241 aaatctccag agattatcag tggcggatg actttcgccc tctgttgcta ttctctgaca 240
243 ttcatgagat ttgccttacaa ggtacaaccc tggaaactggc ttttgtttgc atgccatgt 300
245 acaaacgaag tagctcagct cattcaggga ggacgactta tcaactacga gatgagtaag 360
247 cggccatctg catagcggta caaggaccag ctcttggaaag agacagtgtt ccagccactg 420
249 ctgcagccac agatcatgtc agcatgagta gtcgtgctga agggaaaaca cagaatgcta 480
251 tcttaatgac catgccaaca ttattgaata gcccggatgc cctaaacccca ctctctgctg 540
253 ccttatcaat gctaaacccctt atttgtcttc atcaagagta gttcaaaata tgcaactaat 600
255 ttaataattt tgaatgatgg ttttatctat agcaatctgt agtaatatgt atattatcta 660
257 ttgggatttg tggataaaaa aatctaaggq aacaaaattt tataactaca aqcaacttaaq 720

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259 tactcaaaat tcttgacttt ttctttaatg acaatagtaa accctcagtt ggtcacatgt 780
 261 ctacacataa tttccagtga taacaagtt cgggttttc catatgtaac tcagatctgt 840
 263 aacttaatgg caataaaatgg tttaaatatt tgc 873
 266 <210> SEQ ID NO: 9
 267 <211> LENGTH: 549
 268 <212> TYPE: DNA
 269 <213> ORGANISM: Mus musculus
 271 <400> SEQUENCE: 9
 272 cggcacagcc atggcgggcg cgttggcgcg gaaagcggcg gactatgtcc gaagcaagga 60
 274 tttccggac tacctcatga gtacgcactt ctggggccca gtagccaact ggggtcttcc 120
 276 cattgctgcc atcaatgata tgaaaaagtc tccagagatt atcagtgccg ggtatgacatt 180
 278 tgccctctgt tgctattctt tgacattcat gagatttgcc tacaaggtaa agcctcgaa 240
 280 ctggcttcgt tttgcatgcc acgcaacaaa tgaagtagcc cagctcatcc agggagggcg 300
 282 gcttatcaaa cacgagatga ctgttaactt atgcaataa atgatttaaa tatttgaaga 360
 284 gtagctcaaa atatgcaatt aatttaataa ttatctgca gtaatatgtat tattcatctat 420
 286 tagaatttac ttaatgaaaa actgaagaga acaaatttg taaccactag cacttaagta 480
 288 ctccctgattc ttaacattgt cttaatgac aatagctgag agtttctaaa ccaactctct 540
 290 gtcgcctta 549
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 296 <213> ORGANISM: Artificial Sequence
 298 <220> FEATURE:
 299 <223> OTHER INFORMATION: 5' random arbitrary primer
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 302 caagcgaggt 10
 305 <210> SEQ ID NO: 11
 306 <211> LENGTH: 10
 307 <212> TYPE: DNA
 308 <213> ORGANISM: Artificial Sequence
 310 <220> FEATURE:
 311 <223> OTHER INFORMATION: 5' random arbitrary primer
 313 <400> SEQUENCE: 11
 314 cagttagctg 10
 317 <210> SEQ ID NO: 12
 318 <211> LENGTH: 10
 319 <212> TYPE: DNA
 320 <213> ORGANISM: Artificial Sequence
 322 <220> FEATURE:
 323 <223> OTHER INFORMATION: 5' random arbitrary primer
 325 <400> SEQUENCE: 12
 326 gtcacggaaag 10
 329 <210> SEQ ID NO: 13
 330 <211> LENGTH: 27
 331 <212> TYPE: DNA
 332 <213> ORGANISM: Artificial Sequence
 334 <220> FEATURE:
 335 <223> OTHER INFORMATION: PCR primer AP1
 337 <400> SEQUENCE: 13

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Input Set : A:\2004-08-12 0641-0260P.ST25.txt
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338 ccatccta at acgactca ct ataggc 27
 341 <210> SEQ ID NO: 14
 342 <211> LENGTH: 24
 343 <212> TYPE: DNA
 344 <213> ORGANISM: Artificial Sequence
 346 <220> FEATURE:
 347 <223> OTHER INFORMATION: PCR primer corresponding to SEQ ID NO: 1
 349 <400> SEQUENCE: 14
 350 agccgagagt tggtttgggg attc 24
 353 <210> SEQ ID NO: 15
 354 <211> LENGTH: 22
 355 <212> TYPE: DNA
 356 <213> ORGANISM: Artificial Sequence
 358 <220> FEATURE:
 359 <223> OTHER INFORMATION: Primer I (ARBP cDNA 5' primer)
 361 <400> SEQUENCE: 15
 362 gggatccaac agcacggcca tg 22
 365 <210> SEQ ID NO: 16
 366 <211> LENGTH: 26
 367 <212> TYPE: DNA
 368 <213> ORGANISM: Artificial Sequence
 370 <220> FEATURE:
 371 <223> OTHER INFORMATION: primer II (ARBP cDNA 3' primer)
 373 <400> SEQUENCE: 16
 374 ggaattcatt gataaggcag ccgaga 26
 377 <210> SEQ ID NO: 17
 378 <211> LENGTH: 20
 379 <212> TYPE: DNA
 380 <213> ORGANISM: Artificial Sequence
 382 <220> FEATURE:
 383 <223> OTHER INFORMATION: GAPDH sense primer
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 386 tgctggtgct gagtatgtcg 20
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 391 <212> TYPE: DNA
 392 <213> ORGANISM: Artificial Sequence
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 395 <223> OTHER INFORMATION: GAPDH anti-sense primer
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 398 gcatgtcaga tccacaacgg 20
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 402 <211> LENGTH: 14
 403 <212> TYPE: PRT
 404 <213> ORGANISM: Artificial Sequence
 406 <220> FEATURE:
 407 <223> OTHER INFORMATION: polyclonal antibody against a C-terminal peptide sequence of
 ARBP
 409 <400> SEQUENCE: 19
 411 Gly Arg Leu Ile Asn Tyr Glu Met Ser Lys Arg Pro Ser Ala

VERIFICATION SUMMARY

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